

SEQUENCE LISTING

<110> Reiner, Peter B.
Connop, Bruce P.
Pollard, Michelle

<120> REGULATION OF AMYLOID PRECURSOR PROTEIN EXPRESSION
BY MODIFICATION OF ABC TRANSPORTER EXPRESSION OR ACTIVITY

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<141> 2002-02-08

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 Asp Leu Trp Ala Ala Cys Leu Tyr Arg Ser Cys Leu Leu Leu Gly Ala
 50 55 60
 Thr Ile Gly Val Ala Lys Asn Ser Ala Leu Gly Pro Arg Arg Leu Arg
 65 70 75 80
 Ala Ser Trp Leu Val Ile Thr Leu Val Cys Leu Phe Val Gly Ile Tyr
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 Ala Met Val Lys Leu Leu Leu Phe Ser Glu Val Arg Arg Pro Ile Arg
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 Asp Pro Trp Phe Trp Ala Leu Phe Val Trp Thr Tyr Ile Ser Leu Gly
 115 120 125
 Ala Ser Phe Leu Leu Trp Trp Leu Leu Ser Thr Val Arg Pro Gly Thr
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 690 695 700
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 Gln Gly Thr His Gln Gln Leu Leu Ala Gln Gly Gly Leu Tyr Ala Lys
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Arg Tyr Ala Tyr Tyr Tyr Ser Gly Ile Gly Ala Gly Val Leu Val Ala			
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Ala Tyr Ile Gln Val Ser Phe Trp Cys Leu Ala Ala Gly Arg Gln Ile			
130	135	140	
His Lys Ile Arg Lys Gln Phe Phe His Ala Ile Met Arg Gln Glu Ile			
145	150	155	160
Gly Trp Phe Asp Val His Asp Val Gly Glu Leu Asn Thr Arg Leu Thr			
165	170	175	
Asp Asp Val Ser Lys Ile Asn Glu Gly Ile Gly Asp Lys Ile Gly Met			
180	185	190	
Phe Phe Gln Ser Met Ala Thr Phe Phe Thr Gly Phe Ile Val Gly Phe			
195	200	205	
Thr Arg Gly Trp Lys Leu Thr Leu Val Ile Leu Ala Ile Ser Pro Val			
210	215	220	
Leu Gly Leu Ser Ala Ala Val Trp Ala Lys Ile Leu Ser Ser Phe Thr			
225	230	235	240
Asp Lys Glu Leu Leu Ala Tyr Ala Lys Ala Gly Ala Val Ala Glu Glu			
245	250	255	
Val Leu Ala Ala Ile Arg Thr Val Ile Ala Phe Gly Gly Gln Lys Lys			
260	265	270	
Glu Leu Glu Arg Tyr Asn Lys Asn Leu Glu Glu Ala Lys Arg Ile Gly			
275	280	285	
Ile Lys Lys Ala Ile Thr Ala Asn Ile Ser Ile Gly Ala Ala Phe Leu			
290	295	300	
Leu Ile Tyr Ala Ser Tyr Ala Leu Ala Phe Trp Tyr Gly Thr Thr Leu			
305	310	315	320
Val Leu Ser Gly Glu Tyr Ser Ile Gly Gln Val Leu Thr Val Phe Phe			
325	330	335	

Ser Val Leu Ile Gly Ala Phe Ser Val Gly Gln Ala Ser Pro Ser Ile
 340 345 350
 Glu Ala Phe Ala Asn Ala Arg Gly Ala Ala Tyr Glu Ile Phe Lys Ile
 355 360 365
 Ile Asp Asn Lys Pro Ser Ile Asp Ser Tyr Ser Lys Ser Gly His Lys
 370 375 380
 Pro Asp Asn Ile Lys Gly Asn Leu Glu Phe Arg Asn Val His Phe Ser
 385 390 395 400
 Tyr Pro Ser Arg Lys Glu Val Lys Ile Leu Lys Gly Leu Asn Leu Lys
 405 410 415
 Val Gln Ser Gly Gln Thr Val Ala Leu Val Gly Asn Ser Gly Cys Gly
 420 425 430
 Lys Ser Thr Thr Val Gln Leu Met Gln Arg Leu Tyr Asp Pro Thr Glu
 435 440 445
 Gly Met Val Ser Val Asp Gly Gln Asp Ile Arg Thr Ile Asn Val Arg
 450 455 460
 Phe Leu Arg Glu Ile Ile Gly Val Val Ser Gln Glu Pro Val Leu Phe
 465 470 475 480
 Ala Thr Thr Ile Ala Glu Asn Ile Arg Tyr Gly Arg Glu Asn Val Thr
 485 490 495
 Met Asp Glu Ile Glu Lys Ala Val Lys Glu Ala Asn Ala Tyr Asp Phe
 500 505 510
 Ile Met Lys Leu Pro His Lys Phe Asp Thr Leu Val Gly Glu Arg Gly
 515 520 525
 Ala Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile Ala Arg Ala
 530 535 540
 Leu Val Arg Asn Pro Lys Ile Leu Leu Asp Glu Ala Thr Ser Ala
 545 550 555 560
 Leu Asp Thr Glu Ser Glu Ala Val Val Gln Val Ala Leu Asp Lys Ala
 565 570 575
 Arg Lys Gly Arg Thr Thr Ile Val Ile Ala His Arg Leu Ser Thr Val
 580 585 590
 Arg Asn Ala Asp Val Ile Ala Gly Phe Asp Asp Gly Val Ile Val Glu
 595 600 605
 Lys Gly Asn His Asp Glu Leu Met Lys Glu Lys Gly Ile Tyr Phe Lys
 610 615 620
 Leu Val Thr Met Gln Thr Ala Gly Asn Glu Val Glu Leu Glu Asn Ala
 625 630 635 640
 Ala Asp Glu Ser Lys Ser Glu Ile Asp Ala Leu Glu Met Ser Ser Asn
 645 650 655
 Asp Ser Arg Ser Ser Leu Ile Arg Lys Arg Ser Thr Arg Arg Ser Val
 660 665 670
 Arg Gly Ser Gln Ala Gln Asp Arg Lys Leu Ser Thr Lys Glu Ala Leu
 675 680 685
 Asp Glu Ser Ile Pro Pro Val Ser Phe Trp Arg Ile Met Lys Leu Asn
 690 695 700
 Leu Thr Glu Trp Pro Tyr Phe Val Val Gly Val Phe Cys Ala Ile Ile
 705 710 715 720
 Asn Gly Gly Leu Gln Pro Ala Phe Ala Ile Ile Phe Ser Lys Ile Ile
 725 730 735
 Gly Val Phe Thr Arg Ile Asp Asp Pro Glu Thr Lys Arg Gln Asn Ser
 740 745 750
 Asn Leu Phe Ser Leu Leu Phe Leu Ala Leu Gly Ile Ile Ser Phe Ile
 755 760 765

Thr Phe Phe Leu Gln Gly Phe Thr Phe Gly Lys Ala Gly Glu Ile Leu
 770 775 780
 Thr Lys Arg Leu Arg Tyr Met Val Phe Arg Ser Met Leu Arg Gln Asp
 785 790 795 800
 Val Ser Trp Phe Asp Asp Pro Lys Asn Thr Thr Gly Ala Leu Thr Thr
 805 810 815
 Arg Leu Ala Asn Asp Ala Ala Gln Val Lys Gly Ala Ile Gly Ser Arg
 820 825 830
 Leu Ala Val Ile Thr Gln Asn Ile Ala Asn Leu Gly Thr Gly Ile Ile
 835 840 845
 Ile Ser Phe Ile Tyr Gly Trp Gln Leu Thr Leu Leu Leu Ala Ile
 850 855 860
 Val Pro Ile Ile Ala Ile Ala Gly Val Val Glu Met Lys Met Leu Ser
 865 870 875 880
 Gly Gln Ala Leu Lys Asp Lys Lys Glu Leu Glu Gly Ala Gly Lys Ile
 885 890 895
 Ala Thr Glu Ala Ile Glu Asn Phe Arg Thr Val Val Ser Leu Thr Gln
 900 905 910
 Glu Gln Lys Phe Glu His Met Tyr Ala Gln Ser Leu Gln Val Pro Tyr
 915 920 925
 Arg Asn Ser Leu Arg Lys Ala His Ile Phe Gly Ile Thr Phe Ser Phe
 930 935 940
 Thr Gln Ala Met Met Tyr Phe Ser Tyr Ala Gly Cys Phe Arg Phe Gly
 945 950 955 960
 Ala Tyr Leu Val Ala His Lys Leu Met Ser Phe Glu Asp Val Leu Leu
 965 970 975
 Val Phe Ser Ala Val Val Phe Gly Ala Met Ala Val Gly Gln Val Ser
 980 985 990
 Ser Phe Ala Pro Asp Tyr Ala Lys Ala Lys Ile Ser Ala Ala His Ile
 995 1000 1005
 Ile Met Ile Ile Glu Lys Thr Pro Leu Ile Asp Ser Tyr Ser Thr Glu
 1010 1015 1020
 Gly Leu Met Pro Asn Thr Leu Glu Gly Asn Val Thr Phe Gly Glu Val
 1025 1030 1035 1040
 Val Phe Asn Tyr Pro Thr Arg Pro Asp Ile Pro Val Leu Gln Gly Leu
 1045 1050 1055
 Ser Leu Glu Val Lys Lys Gly Gln Thr Leu Ala Leu Val Gly Ser Ser
 1060 1065 1070
 Gly Cys Gly Lys Ser Thr Val Val Gln Leu Leu Glu Arg Phe Tyr Asp
 1075 1080 1085
 Pro Leu Ala Gly Lys Val Leu Leu Asp Gly Lys Glu Ile Lys Arg Leu
 1090 1095 1100
 Asn Val Gln Trp Leu Arg Ala His Leu Gly Ile Val Ser Gln Glu Pro
 1105 1110 1115 1120
 Ile Leu Phe Asp Cys Ser Ile Ala Glu Asn Ile Ala Tyr Gly Asp Asn
 1125 1130 1135
 Ser Arg Val Val Ser Gln Glu Glu Ile Val Arg Ala Ala Lys Glu Ala
 1140 1145 1150
 Asn Ile His Ala Phe Ile Glu Ser Leu Pro Asn Lys Tyr Ser Thr Lys
 1155 1160 1165
 Val Gly Asp Lys Gly Thr Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile
 1170 1175 1180
 Ala Ile Ala Arg Ala Leu Val Arg Gln Pro His Ile Leu Leu Leu Asp
 1185 1190 1195 1200

Glu Ala Thr Ser Ala Leu Asp Thr Glu Ser Glu Lys Val Val Gln Glu
 1205 1210 1215
 Ala Leu Asp Lys Ala Arg Glu Gly Arg Thr Cys Ile Val Ile Ala His
 1220 1225 1230
 Arg Leu Ser Thr Ile Gln Asn Ala Asp Leu Ile Val Val Phe Gln Asn
 1235 1240 1245
 Gly Arg Val Lys Glu His Gly Thr His Gln Gln Leu Leu Ala Gln Lys
 1250 1255 1260
 Gly Ile Tyr Phe Ser Met Val Ser Val Gln Ala Gly Thr Lys Arg Gln
 1265 1270 1275 1280

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 <213> Homo sapiens

<220>
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 <223> Xaa = Any Amino Acid

<221> VARIANT
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 <223> Xaa = Any Amino Acid

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 Leu Val His Leu Met Thr Ser Asn Pro Lys Ile Leu Tyr Xaa Pro Ala
 20 25 30
 Gly Ser Glu Val Asp Arg Val Ile Leu Lys Ala Asn Glu Thr Phe Ala
 35 40 45
 Phe Val Gly Asn Val Thr His Tyr Ala Gln Val Trp Leu Asn Ile Ser
 50 55 60
 Ala Glu Ile Arg Ser Xaa Leu Glu Gin Gly Arg Leu Gln Gln His Leu
 65 70 75 80
 Arg Trp Leu Gln Gln Tyr Val Ala Glu Leu Arg Pro His Pro Glu Ala
 85 90 95
 Leu Asn Leu Ser Leu Asp Glu Leu Pro Pro Ala Leu Arg Gln Asp Asn
 100 105 110
 Phe Ser Leu Pro Ser Gly Met Ala Leu Leu Gln Gln Leu Asp Thr Ile
 115 120 125
 Asp Asn Ala Pro Cys Gly Trp Ile Gln Phe Met Ser Lys Val Ser Val
 130 135 140
 Asp Ile Phe Lys Gly Phe Pro Asp Glu Glu Ser Ile Val Asn Tyr Thr
 145 150 155 160
 Leu Asn Gln Ala Tyr Gln Asp Asn Val Thr Val Phe Ala Gly Val Ile
 165 170 175
 Phe Gln Thr Arg Lys Asp Gly Ser Leu Pro Pro His Val His Tyr Lys
 180 185 190
 Ile Arg Gln Asn Ser Ser Phe Thr Glu Lys Thr Asn Glu Ile Arg Arg
 195 200 205
 Ala Tyr Trp Arg Pro Gly Pro Asn Thr Gly Gly Arg Phe Tyr Phe Leu

210	215	220													
Tyr	Gly	Phe	Val	Trp	Ile	Gln	Asp	Met	Met	Glu	Arg	Ala	Ile	Ile	Asp
225				230				235			240				
Thr	Phe	Val	Gly	His	Asp	Val	Val	Glu	Pro	Gly	Ser	Tyr	Val	Gln	Met
							245		250				255		
Phe	Pro	Tyr	Pro	Cys	Tyr	Thr	Arg	Asp	Asp	Phe	Leu	Phe	Val	Ile	Glu
						260		265			270				
His	Met	Met	Pro	Leu	Cys	Met	Xaa	Ile	Ser	Trp	Val	Tyr	Ser	Val	Ala
						275		280			285				
Met	Thr	Ile	Gln	His	Ile	Val	Ala	Glu	Lys	Glu	His	Arg	Leu	Lys	Glu
						290		295			300				
Val	Met	Lys	Thr	Met	Gly	Leu	Asn	Asn	Ala	Val	His	Trp	Val	Ala	Trp
305					310				315			320			
Phe	Ile	Thr	Gly	Phe	Val	Gln	Leu	Ser	Ile	Ser	Val	Thr	Ala	Leu	Thr
						325		330			335				
Ala	Ile	Leu	Lys	Tyr	Gly	Gln	Val	Leu	Met	His	Ser	His	Val	Val	Ile
						340		345			350				
Ile	Trp	Leu	Phe	Leu	Ala	Val	Tyr	Ala	Val	Ala	Thr	Ile	Met	Phe	Cys
						355		360			365				
Phe	Leu	Val	Ser	Val	Leu	Tyr	Ser	Lys	Ala	Lys	Leu	Ala	Ser	Ala	Gly
						370		375			380				
Gly	Ile	Ile	Tyr	Phe	Leu	Ser	Tyr	Val	Pro	Tyr	Met	Tyr	Val	Ala	Ile
385					390				395			400			
Arg	Glu	Glu	Val	Ala	His	Asp	Lys	Ile	Thr	Ala	Phe	Glu	Lys	Cys	Ile
						405		410			415				
Ala	Ser	Leu	Met	Ser	Thr	Thr	Ala	Phe	Gly	Leu	Gly	Ser	Lys	Tyr	Phe
						420		425			430				
Ala	Leu	Tyr	Glu	Val	Ala	Gly	Val	Gly	Ile	Gln	Trp	His	Thr	Phe	Ser
						435		440			445				
Gln	Ser	Pro	Val	Glu	Gly	Asp	Asp	Phe	Asn	Leu	Leu	Leu	Ala	Val	Thr
						450		455			460				
Met	Leu	Met	Val	Asp	Ala	Val	Val	Tyr	Gly	Ile	Leu	Xaa	Trp	Tyr	Ile
465						470			475			480			
Glu	Ala	Val	His	Pro	Gly	Met	Tyr	Gly	Leu	Pro	Arg	Pro	Trp	Tyr	Phe
						485		490			495				
Pro	Leu	Gln	Lys	Ser	Tyr	Trp	Leu	Gly	Ser	Gly	Arg	Thr	Glu	Ala	Trp
						500		505			510				
Glu	Trp	Ser	Trp	Pro	Trp	Ala	Arg	Thr	Pro	Arg	Leu	Ser	Val	Met	Glu
						515		520			525				
Glu	Asp	Gln	Ala	Cys	Ala	Met	Glu	Ser	Arg	Arg	Phe	Glu	Glu	Thr	Arg
						530		535			540				
Gly	Met	Glu	Glu	Glu	Pro	Thr	His	Leu	Pro	Leu	Val	Val	Xaa	Val	Asp
545						550			555			560			
Lys	Leu	Thr	Lys	Val	Tyr	Lys	Asp	Asp	Lys	Lys	Leu	Ala	Leu	Asn	Lys
						565		570			575				
Leu	Ser	Leu	Asn	Leu	Tyr	Glu	Asn	Gln	Gly	Val	Ser	Phe	Leu	Gly	His
						580		585			590				
Asn	Gly	Ala	Gly	Lys	Thr	Thr	Met	Ser	Ile	Leu	Thr	Gly	Leu	Phe	
						595		600			605				
Pro	Pro	Thr	Ser	Gly	Ser	Ala	Thr	Ile	Tyr	Gly	His	Asp	Ile	Arg	Thr
						610		615			620				
Glu	Met	Asp	Glu	Ile	Arg	Lys	Asn	Gly	His	Val	Pro	Gln	His	Asn	Val
625						630			635			640			
Leu	Phe	Asp	Arg	Leu	Thr	Val	Glu	Glu	His	Leu	Trp	Phe	Tyr	Ser	Arg

645	650	655
Leu Lys Ser Met Ala Gln Glu Glu Ile Pro Arg Glu Met Asp Lys Met		
660	665	670
Ile Glu Asp Leu Glu Leu Ser Asn Lys Arg His Ser Leu Val Gln Thr		
675	680	685
Leu Ser Gly Gly Met Lys Arg Lys Val Ser Val Ala Ile Ala Phe Val		
690	695	700
Gly Gly Ser Arg Ala Ile Ile Leu Asp Glu Pro Thr Ala Gly Val Asp		
705	710	715
Pro Tyr Ala Arg Arg Ala Ile Trp Asp Leu Ile Leu Lys Tyr Lys Pro		
725	730	735
Gly Arg Thr Ile Leu Leu Ser Thr His His Met Asp Glu Ala Asp Leu		
740	745	750
Leu Gly Asp Arg Ile Ala Ile Ile Ser His Gly Lys Leu Lys Cys Cys		
755	760	765
Gly Ser Pro Leu Phe Leu Lys Gly Thr Tyr Gly Asp Gly Tyr Arg Leu		
770	775	780
Thr Leu Val Lys Arg Pro Ala Glu Pro Gly Gly Pro Gln Glu Pro Gly		
785	790	795
Leu Ala Ser Ser Pro Pro Gly Arg Ala Pro Leu Ser Ser Cys Ser Glu		
805	810	815
Leu Gln Val Ser Gln Phe Ile Arg Lys His Val Ala Ser Cys Leu Leu		
820	825	830
Val Ser Asp Thr Ser Thr Glu Leu Ser Tyr Ile Leu Pro Ser Glu Ala		
835	840	845
Ala Lys Lys Gly Ala Phe Glu Arg Leu Phe Gln His Leu Glu Arg Ser		
850	855	860
Leu Asp Ala Leu His Leu Ser Ser Phe Gly Leu Met Asp Thr Thr Leu		
865	870	875
Glu Glu Val Phe Leu Lys Val Ser Gly Gly Asp Gln Ser Leu Glu Asn		
885	890	895
Ser Gly Ala Asp Val Lys Glu Ser Arg Lys Asp Val Leu Pro Gly Ala		
900	905	910
Glu Gly His Ala Ser Gly Glu Gly His Ala Gly Asn Leu Ala Arg Cys		
915	920	925
Ser Glu Leu Thr Gln Ser Gln Ala Ser Leu Gln Ser Ala Ser Ser Val		
930	935	940
Gly Ser Ala Leu Gly Asp Glu Gly Ala Gly Tyr Thr Asp Val Tyr Gly		
945	950	955
Asp Tyr Pro Pro Leu Phe Asp Asn Pro Gln Asp Pro Asp Asn Val Ser		
965	970	975
Leu Gln Glu Val Glu Ala Glu Ala Leu Ser Arg Val Gly Gln Gly Ser		
980	985	990
Arg Lys Leu Asp Gly Gly Trp Leu Lys Val Arg Gln Phe His Gly Leu		
995	1000	1005
Leu Val Lys Arg Phe His Cys Ala Arg Arg Asn Ser Lys Ala Leu Phe		
1010	1015	1020
Ser Gln Ile Leu Leu Pro Ala Phe Phe Val Cys Val Ala Met Thr Val		
1025	1030	1035
Ala Leu Ser Val Pro Glu Ile Gly Asp Leu Pro Pro Leu Val Leu Ser		
1045	1050	1055
Pro Ser Gln Tyr His Asn Tyr Thr Gln Pro Arg Gly Asn Phe Ile Pro		
1060	1065	1070
Tyr Ala Asn Glu Glu Arg Arg Glu Tyr Arg Leu Arg Leu Ser Pro Asp		

1075	1080	1085
Ala Ser Pro Gln Gln Leu Val	Ser Thr Phe Arg Leu Pro Ser Gly Val	
1090	1095	1100
Gly Ala Thr Cys Val Leu Lys Ser Pro Ala Asn Gly Ser Leu Gly Pro		
1105	1110	1115
Thr Leu Asn Leu Ser Ser Gly Glu Ser Arg Leu Leu Ala Ala Arg Phe		
1125	1130	1135
Phe Asp Ser Met Cys Leu Glu Ser Phe Thr Gln Gly Leu Pro Leu Ser		
1140	1145	1150
Asn Phe Val Pro Pro Pro Ser Pro Ala Pro Ser Asp Ser Pro Ala		
1155	1160	1165
Ser Pro Asp Glu Asp Leu Gln Ala Trp Asn Val Ser Leu Pro Pro Thr		
1170	1175	1180
Ala Gly Gln Glu Met Trp Thr Ser Ala Pro Ser Leu Pro Arg Leu Val		
1185	1190	1195
Arg Glu Pro Val Arg Cys Thr Cys Ser Ala Gln Gly Thr Gly Phe Ser		
1205	1210	1215
Cys Pro Asn Ser Val Gly Gly His Pro Pro Gln Met Arg Val Val Thr		
1220	1225	1230
Gly Asp Ile Leu Thr Asp Ile Thr Gly His Asn Val Ser Glu Tyr Leu		
1235	1240	1245
Leu Phe Thr Ser Asp Arg Phe Arg Leu His Arg Tyr Gly Ala Ile Thr		
1250	1255	1260
Phe Gly Asn Val Leu Lys Ser Ile Pro Ala Ser Phe Gly Thr Arg Ala		
1265	1270	1275
Pro Pro Met Val Arg Lys Ile Arg Cys Ala Arg Ala Ala Gln Val Phe		
1285	1290	1295
Tyr Asn Asn Lys Gly Tyr His Ser Met Pro Thr Tyr Leu Asn Ser Leu		
1300	1305	1310
Asn Asn Ala Ile Leu Arg Ala Asn Leu Pro Lys Ser Lys Gly Asn Pro		
1315	1320	1325
Ala Ala Tyr Gly Ile Thr Val Thr Asn His Pro Met Asn Lys Thr Ser		
1330	1335	1340
Ala Ser Leu Ser Leu Asp Tyr Leu Leu Gln Gly Thr Asp Val Val Ile		
1345	1350	1355
Ala Ile Phe Ile Ile Val Ala Met Ser Phe Val Pro Ala Ser Phe Val		
1365	1370	1375
Val Phe Leu Val Ala Glu Lys Ser Thr Lys Ala Lys His Leu Gln Phe		
1380	1385	1390
Val Ser Gly Cys Asn Pro Ile Ile Tyr Trp Leu Ala Asn Tyr Val Trp		
1395	1400	1405
Asp Met Leu Asn Tyr Leu Val Pro Ala Thr Cys Cys Val Ile Ile Leu		
1410	1415	1420
Phe Val Phe Asp Leu Pro Ala Tyr Thr Ser Pro Thr Asn Phe Pro Ala		
1425	1430	1435
Val Leu Ser Leu Phe Leu Leu Tyr Gly Trp Ser Ile Thr Pro Ile Met		
1445	1450	1455
Tyr Pro Ala Ser Phe Trp Phe Glu Val Pro Ser Ser Ala Tyr Xaa Phe		
1460	1465	1470
Leu Ile Val Ile Asn Leu Phe Ile Gly Ile Thr Ala Thr Val Ala Thr		
1475	1480	1485
Phe Leu Leu Gln Leu Phe Glu His Asp Lys Asp Leu Lys Val Val Asn		
1490	1495	1500
Ser Tyr Leu Lys Ser Cys Phe Leu Ile Phe Pro Asn Tyr Asn Leu Gly		

1505	1510	1515	1520												
His	Gly	Leu	Met	Glu	Met	Ala	Tyr	Asn	Glu	Tyr	Ile	Asn	Glu	Tyr	Tyr
				1525				1530							1535
Ala	Lys	Ile	Gly	Gln	Phe	Asp	Lys	Met	Lys	Ser	Pro	Phe	Glu	Trp	Asp
				1540				1545							1550
Ile	Val	Thr	Arg	Gly	Leu	Val	Ala	Met	Ala	Val	Glu	Gly	Val	Val	Gly
				1555				1560							1565
Phe	Leu	Leu	Thr	Ile	Met	Cys	Gln	Tyr	Asn	Phe	Leu	Arg	Arg	Pro	Gln
				1570				1575							1580
Arg	Met	Pro	Val	Ser	Thr	Lys	Pro	Val	Glu	Asp	Asp	Val	Asp	Val	Ala
				1585				1590							1600
Ser	Glu	Arg	Gln	Arg	Val	Leu	Arg	Gly	Asp	Ala	Asp	Asn	Asp	Met	Val
				1605				1610							1615
Lys	Ile	Glu	Asn	Leu	Thr	Lys	Val	Tyr	Lys	Ser	Arg	Lys	Ile	Gly	Arg
				1620				1625							1630
Ile	Leu	Ala	Val	Asp	Arg	Leu	Cys	Leu	Gly	Val	Arg	Pro	Gly	Glu	Cys
				1635				1640							1645
Phe	Gly	Xaa	Leu	Gly	Val	Asn	Gly	Ala	Gly	Lys	Thr	Ser	Thr	Phe	Lys
				1650				1655							1660
Met	Leu	Thr	Gly	Asp	Glu	Ser	Thr	Thr	Gly	Gly	Glu	Ala	Phe	Val	Asn
				1665				1670							1680
Gly	His	Ser	Val	Leu	Lys	Glu	Leu	Xaa	Gln	Val	Gln	Gln	Ser	Leu	Gly
				1685				1690							1695
Tyr	Cys	Pro	Gln	Cys	Asp	Ala	Leu	Phe	Asp	Glu	Leu	Thr	Ala	Arg	Glu
				1700				1705							1710
His	Leu	Gln	Leu	Tyr	Thr	Arg	Leu	Arg	Gly	Ile	Xaa	Trp	Lys	Asp	Glu
				1715				1720							1725
Ala	Arg	Val	Val	Lys	Trp	Ala	Leu	Glu	Lys	Leu	Glu	Leu	Thr	Lys	Tyr
				1730				1735							1740
Ala	Asp	Lys	Pro	Ala	Gly	Thr	Tyr	Ser	Gly	Gly	Asn	Lys	Arg	Lys	Leu
				1745				1750							1760
Ser	Thr	Ala	Ile	Ala	Leu	Ile	Gly	Tyr	Pro	Ala	Phe	Ile	Phe	Leu	Asp
				1765				1770							1775
Glu	Pro	Thr	Thr	Gly	Met	Asp	Pro	Lys	Ala	Arg	Arg	Phe	Leu	Trp	Asn
				1780				1785							1790
Leu	Ile	Leu	Asp	Leu	Ile	Lys	Thr	Gly	Arg	Ser	Val	Val	Leu	Thr	Ser
				1795				1800							1805
His	Ser	Met	Glu	Glu	Cys	Glu	Ala	Leu	Cys	Thr	Arg	Leu	Ala	Ile	Met
				1810				1815							1820
Val	Asn	Gly	Arg	Leu	Arg	Cys	Leu	Gly	Ser	Ile	Gln	His	Leu	Lys	Asn
				1825				1830							1840
Arg	Phe	Gly	Asp	Gly	Tyr	Met	Ile	Thr	Val	Arg	Thr	Lys	Ser	Ser	Gln
				1845				1850							1855
Ser	Val	Lys	Asp	Val	Val	Arg	Phe	Phe	Asn	Arg	Asn	Phe	Pro	Glu	Ala
				1860				1865							1870
Met	Leu	Lys	Glu	Arg	His	His	Thr	Lys	Val	Gln	Tyr	Gln	Leu	Lys	Ser
				1875				1880							1885
Glu	His	Ile	Ser	Leu	Ala	Gln	Val	Phe	Ser	Lys	Met	Glu	Gln	Val	Ser
				1890				1895							1900
Gly	Val	Leu	Gly	Ile	Glu	Asp	Tyr	Ser	Val	Ser	Gln	Thr	Leu	Asp	
				1905				1910							1920
Asn	Val	Phe	Val	Asn	Phe	Ala	Lys	Lys	Gln	Ser	Asp	Asn	Leu	Glu	Gln
				1925				1930							1935
Gln	Glu	Thr	Glu	Pro	Pro	Ser	Ala	Leu	Gln	Ser	Pro	Leu	Gly	Cys	Leu

1940	1945	1950
Leu Ser Leu Leu Arg Pro Arg Ser Ala Pro Thr Glu Leu Arg Ala Leu		
1955	1960	1965
Val Ala Asp Glu Pro Glu Asp Leu Asp Thr Glu Asp Glu Gly Leu Ile		
1970	1975	1980
Ser Phe Glu Glu Glu Arg Ala Gln Leu Ser Phe Asn Thr Asp Thr Leu		
1985	1990	1995
Cys		2000

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<211> 646
<212> PRT
<213> Homo sapiens

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Ala Val Ala Met Ala Val Thr Leu Glu Asp Gly Ala Glu Pro Pro Val
20 25 30
Leu Thr Thr His Leu Lys Lys Val Glu Asn His Ile Thr Glu Ala Gln
35 40 45
Arg Phe Ser His Leu Pro Lys Arg Ser Ala Val Asp Ile Glu Phe Val
50 55 60
Glu Leu Ser Tyr Ser Val Arg Glu Gly Pro Cys Trp Arg Lys Arg Gly
65 70 75 80
Tyr Lys Thr Leu Leu Lys Cys Leu Ser Gly Lys Phe Cys Arg Arg Glu
85 90 95
Leu Ile Gly Ile Met Gly Pro Ser Gly Ala Gly Lys Ser Thr Phe Met
100 105 110
Asn Ile Leu Ala Gly Tyr Arg Glu Ser Gly Met Lys Gly Gln Ile Leu
115 120 125
Val Asn Gly Arg Pro Arg Glu Leu Arg Thr Phe Arg Lys Met Ser Cys
130 135 140
Tyr Ile Met Gln Asp Asp Met Leu Leu Pro His Leu Thr Val Leu Glu
145 150 155 160
Ala Met Met Val Ser Ala Asn Leu Asn Leu Thr Glu Asn Pro Asp Val
165 170 175
Lys Asn Asp Leu Val Thr Glu Ile Leu Thr Ala Leu Gly Leu Met Ser
180 185 190
Cys Ser His Thr Arg Thr Ala Leu Leu Ser Gly Gly Gln Arg Lys Arg
195 200 205
Leu Ala Ile Ala Leu Glu Leu Val Asn Asn Pro Pro Val Met Phe Phe
210 215 220
Asp Glu Pro Thr Ser Gly Leu Asp Ser Ala Ser Cys Phe Gln Val Val
225 230 235 240
Ser Leu Met Lys Ser Leu Ala Gln Gly Gly Arg Thr Ile Ile Cys Thr
245 250 255
Ile His Gln Pro Ser Ala Lys Leu Phe Glu Met Phe Asp Lys Leu Tyr
260 265 270
Ile Leu Ser Gln Gly Gln Cys Ile Phe Lys Gly Val Val Thr Asn Leu
275 280 285
Ile Pro Tyr Leu Lys Gly Leu Gly Leu His Cys Pro Thr Tyr His Asn

290	295	300
Pro Ala Asp Phe Ile Ile Glu Val Ala Ser Gly	Glu Tyr Gly Asp Leu	
305	310	315
Asn Pro Met Leu Phe Arg Ala Val Gln Asn Gly	Leu Cys Ala Met Ala	320
325	330	335
Glu Lys Lys Ser Ser Pro Glu Lys Asn Glu Val	Pro Ala Pro Cys Pro	
340	345	350
Pro Cys Pro Pro Glu Val Asp Pro Ile Glu Ser His	Thr Phe Ala Thr	
355	360	365
Ser Thr Leu Thr Gln Phe Cys Ile Leu Phe Lys Arg	Thr Phe Leu Ser	
370	375	380
Ile Leu Arg Asp Thr Val Leu Thr His Leu Arg Phe	Met Ser His Val	
385	390	395
Val Ile Gly Val Leu Ile Gly Leu Leu Tyr Leu His	Ile Gly Asp Asp	400
405	410	415
Ala Ser Lys Val Phe Asn Asn Thr Gly Cys Leu Phe	Phe Ser Met Leu	
420	425	430
Phe Leu Met Phe Ala Ala Leu Met Pro Thr Val Leu	Thr Phe Pro Leu	
435	440	445
Glu Met Ala Val Phe Met Arg Glu His Leu Asn Tyr	Trp Tyr Ser Leu	
450	455	460
Lys Ala Tyr Tyr Leu Ala Lys Thr Met Ala Asp Val	Pro Phe Gln Val	
465	470	475
Val Cys Pro Val Val Tyr Cys Ser Ile Val Tyr Trp	Met Thr Gly Gln	
485	490	495
Pro Ala Glu Thr Ser Arg Phe Leu Leu Phe Ser Ala	Leu Ala Thr Ala	
500	505	510
Thr Ala Leu Val Ala Gln Ser Leu Gly Leu Leu	Ile Gly Ala Ala Ser	
515	520	525
Asn Ser Leu Gln Val Ala Thr Phe Val Gly Pro Val	Thr Ala Ile Pro	
530	535	540
Val Leu Leu Phe Ser Gly Phe Phe Val Ser Phe	Lys Thr Ile Pro Thr	
545	550	555
Tyr Leu Gln Trp Ser Ser Tyr Leu Ser Tyr Val Arg	Tyr Gly Phe Glu	
565	570	575
Gly Val Ile Leu Thr Ile Tyr Gly Met Glu Arg Gly	Asp Leu Thr Cys	
580	585	590
Leu Glu Glu Arg Cys Pro Phe Arg Glu Pro Gln Ser	Ile Leu Arg Ala	
595	600	605
Leu Asp Val Glu Asp Ala Lys Leu Tyr Met Asp Phe	Leu Val Leu Gly	
610	615	620
Ile Phe Phe Leu Ala Leu Arg Leu Leu Ala Tyr	Leu Val Leu Arg Tyr	
625	630	635
Arg Val Lys Ser Glu Arg		640
	645	

<210> 10

<211> 638

<212> PRT

<213> Homo sapiens

<400> 10

Met Glu Ala Thr Glu Thr Asp Leu Leu Asn Gly His Leu Lys Lys Val

1	5	10	15
Asp Asn Asn Leu Thr Glu Ala Gln Arg Phe Ser Ser Leu Pro Arg Arg			
20	25	30	
Ala Ala Val Asn Ile Glu Phe Arg Asp Leu Ser Tyr Ser Val Pro Glu			
35	40	45	
Gly Pro Trp Trp Arg Lys Lys Gly Tyr Lys Thr Leu Leu Lys Gly Ile			
50	55	60	
Ser Gly Lys Phe Asn Ser Gly Glu Leu Val Ala Ile Met Gly Pro Ser			
65	70	75	80
Gly Ala Gly Lys Ser Thr Leu Met Asn Ile Leu Ala Gly Tyr Arg Glu			
85	90	95	
Thr Gly Met Lys Gly Ala Val Leu Ile Asn Gly Leu Pro Arg Asp Leu			
100	105	110	
Arg Cys Phe Arg Lys Val Ser Cys Tyr Ile Met Gln Asp Asp Met Leu			
115	120	125	
Leu Pro His Leu Thr Val Gln Glu Ala Met Met Val Ser Ala His Leu			
130	135	140	
Lys Leu Gln Glu Lys Asp Glu Gly Arg Arg Glu Met Val Lys Glu Ile			
145	150	155	160
Leu Thr Ala Leu Gly Leu Leu Ser Cys Ala Asn Thr Arg Thr Gly Ser			
165	170	175	
Leu Ser Gly Gly Gln Arg Lys Arg Leu Ala Ile Ala Leu Glu Leu Val			
180	185	190	
Asn Asn Pro Pro Val Met Phe Phe Asp Glu Pro Thr Ser Gly Leu Asp			
195	200	205	
Ser Ala Ser Cys Phe Gln Val Val Ser Leu Met Lys Gly Leu Ala Gln			
210	215	220	
Gly Gly Arg Ser Ile Ile Cys Thr Ile His Gln Pro Ser Ala Lys Leu			
225	230	235	240
Phe Glu Leu Phe Asp Gln Leu Tyr Val Leu Ser Gln Gly Gln Cys Val			
245	250	255	
Tyr Arg Gly Lys Val Cys Asn Leu Val Pro Tyr Leu Arg Asp Leu Gly			
260	265	270	
Leu Asn Cys Pro Thr Tyr His Asn Pro Ala Asp Phe Val Met Glu Val			
275	280	285	
Ala Ser Gly Glu Tyr Gly Asp Gln Asn Ser Arg Leu Val Arg Ala Val			
290	295	300	
Arg Glu Gly Met Cys Asp Ser Asp His Lys Arg Asp Leu Gly Gly Asp			
305	310	315	320
Ala Glu Val Asn Pro Phe Leu Trp His Arg Pro Ser Glu Glu Val Lys			
325	330	335	
Gln Thr Lys Arg Leu Lys Gly Leu Arg Lys Asp Ser Ser Ser Met Glu			
340	345	350	
Gly Cys His Ser Phe Ser Ala Ser Cys Leu Thr Gln Phe Cys Ile Leu			
355	360	365	
Phe Lys Arg Thr Phe Leu Ser Ile Met Arg Asp Ser Val Leu Thr His			
370	375	380	
Leu Arg Ile Thr Ser His Ile Gly Ile Gly Leu Leu Ile Gly Leu Leu			
385	390	395	400
Tyr Leu Gly Ile Gly Asn Glu Ala Lys Lys Val Leu Ser Asn Ser Gly			
405	410	415	
Phe Leu Phe Phe Ser Met Leu Phe Leu Met Phe Ala Ala Leu Met Pro			
420	425	430	
Thr Val Leu Thr Phe Pro Leu Glu Met Gly Val Phe Leu Arg Glu His			

435	440	445
Leu Asn Tyr Trp Tyr Ser Leu Lys Ala Tyr Tyr	Leu Ala Lys Thr Met	
450	455	460
Ala Asp Val Pro Phe Gln Ile Met Phe Pro Val Ala Tyr Cys Ser Ile		
465	470	475
Val Tyr Trp Met Thr Ser Gln Pro Ser Asp Ala Val Ala Phe Val Leu		480
485	490	495
Phe Ala Ala Leu Gly Thr Met Thr Ser Leu Val Ala Gln Ser Leu Gly		
500	505	510
Leu Leu Ile Gly Ala Ala Ser Thr Ser Leu Gln Val Ala Thr Phe Val		
515	520	525
Gly Pro Val Thr Ala Ile Pro Val Leu Leu Phe Ser Gly Phe Phe Val		
530	535	540
Ser Phe Asp Thr Ile Pro Thr Tyr Leu Gln Trp Met Ser Tyr Ile Ser		
545	550	555
Tyr Val Arg Tyr Gly Phe Glu Gly Val Ile Leu Ser Ile Tyr Gly Leu		560
565	570	575
Asp Arg Glu Asp Leu His Cys Asp Ile Asp Glu Thr Cys His Phe Gln		
580	585	590
Lys Ser Glu Ala Ile Leu Arg Glu Leu Asp Val Glu Asn Ala Lys Leu		
595	600	605
Tyr Leu Asp Phe Ile Val Leu Gly Ile Phe Phe Ile Ser Leu Arg Leu		
610	615	620
Ile Ala Tyr Phe Val Leu Arg Tyr Lys Ile Arg Ala Glu Arg		
625	630	635